

Figure 1.

Alignment of fungal lipolytic enzyme sequences

	1					50	
SEQ ID NO: 1	EVSQDLFNQF	NLFAQYSAAA	YCG	KNNDAPAGTN	33
SEQ ID NO: 2	AV	GVTTFDFS NF	KFYIQHGAAA	YC. .NSEAAAGSK	33
SEQ ID NO: 3	SSSSTQDYRI	ASEAEIKAHT	FYTALSANA.	YCR	TVIPG.....	
SEQ ID NO: 4	.SSSTQDYRI	ASEAEIKAHT	FYTALSANA.	YCR	TVIPG.....	
SEQ ID NO: 5	.SIDGGIRA	ATSQEI NELT	YYTTL SANS.	YCR	TVIPG.....	
SEQ ID NO: 6	.SASDGGKV	AATTAQIQEF	TKYAGIAATA	YCR	SVVPG.....	
SEQ ID NO: 7TAGQAL	AASTQ. GISE	DLYNRL. VEM	ATISQAAYAD	LCNIPST...		
SEQ ID NO: 8TAGHAL	AASTQ. GISE	DLYSRL. VEM	ATISQAAYAD	LCNIPST...		
SEQ ID NO: 9	TVTQDL SNF	RFYLQHADAA	YC. .NFNTAVGKP		
SEQ ID NO: 10	DIPTTQLEDF	KFWVQYAAAT	YCP	NNYVAKDGEK	
SEQ ID NO: 11	DVSTSEL DQF	EFWVQYAAAS	YYE	ADYTAQVGD	
SEQ ID NO: 12	SVSTSTLDEL	QLFAQWSAAA	YCS	NNID. SKDSN	
SEQ ID NO: 13	SVSTSTLDEL	QLFSQWSAAA	YCS	NNID. SDDSN	
SEQ ID NO: 14	DVSSSL LNNL	DLFAQYSAAA	YCD	ENLN. STGTK	
	51					100	
SEQ ID NO: 1	ITCTGNACPE	VEKADATFLY	SFE. DSGVGD	VTGFLALDNT	NKLIVLSFRG		82
SEQ ID NO: 2	ITCSNNGCPT	VQNGATIVT	SF. .VGSKTG	IGGYVATDSA	RKEIVVSFRG		81
SEQ ID NO: 3	GRWSCPHCGV	AS. .NLQITK	TFST. .LITD	TNVLVAVGEK	EKTIYVVFGR		
SEQ ID NO: 4	GQWSCPHCDV	AP. .NLNITK	TFTT. .LITD	TNVLVAVGEN	EKTIYVVFGR		
SEQ ID NO: 5	ATWDCIHCD	TE. .DLKIIK	TWST. .LIYD	TNAMVARGDS	EKTIYVVFGR		
SEQ ID NO: 6	NKWDCVQCQK	WVP. DGKIIT	TFTS. .LLSD	TNGYVLRSDK	QKTIYLVFRG		
SEQ ID NO: 7	IIK	GEKIYNAQTD	INGWILRDDT	SKEIITVFRG	
SEQ ID NO: 8	IIK	GEKIYNSQTD	INGWILRDDT	SKEIITVFRG	
SEQ ID NO: 9	VHCSAGNCPD	IEKDAAIVVG	SV. .VGTKTG	IGAYVATDNA	RKEIVVSFRG		
SEQ ID NO: 10	LNC SVGNCPD	VEAAGSTVKL	SFS. DDTITD	TAGFVAVDNT	NKAI VVAFRG		
SEQ ID NO: 11	LSCSKGNCPE	VEATGATVSY	DFS. DSTITD	TAGYIAVDHT	NSAVVLAFRG		
SEQ ID NO: 12	LTCTANACPS	VEEASTMMLL	EFDLTNDFFG	TAGFLAADNT	NKRLVVAFRG		
SEQ ID NO: 13	VTCTADACPS	VEEASTKMLL	EFDLTNNFFG	TAGFLAADNT	NKRLVVAFRG		
SEQ ID NO: 14	LTCSVGNCPL	VEAASTQSLD	EFNESSSYGN	PAGYLAADET	NKLLVLVSFRG		
	101					150	
SEQ ID NO: 1	SRSIENWIGN	LNFDLKEIND	I. .CSGCRGH	DGFTSSWRSV	ADTLRQKVED		130
SEQ ID NO: 2	SINIRNWL TN	LDFG. QEDCS	L. .VSGCGVH	SGFQRAWNEI	SSQATAAVAS		128
SEQ ID NO: 3	TSSIRNAIAD	IVFVPVNYPP	V. .NGAKVH	KGFLDSYNEV	QDKLVAEVKA		
SEQ ID NO: 4	TSSIRNAIAD	IVFVPVNYPP	V. .NGAKVH	KGFLDSYNEV	QDKLVAEVKA		
SEQ ID NO: 5	SSSIRNWIAD	LTFVPVSYP	V. .SGTKVH	KGFLDSYGEV	QNELVATVLD		
SEQ ID NO: 6	TNSFRSAITD	IVFNFSYK	V. .KGAKVH	AGFLSSYEQV	VNDYFPPVQ		
SEQ ID NO: 7	TGSDTNLQLD	TNYTLTPFDT	LPQCNDCEVH	GGYYIGWISV	QDQVESLVKQ		
SEQ ID NO: 8	TGSDTNLQLD	TNYTLTPFDT	LPQCNSCEVH	GGYYIGWISV	QDQVESLVKQ		
SEQ ID NO: 9	SINVRNWI TN	FNFG. QKTC	L. .VAGCGVH	TGFLDAWEEV	AANVKA AVSA		
SEQ ID NO: 10	SYSIRN WVTD	ATFP. QTDPG	L. .CDGCKAE	LGFWTAWKV	RDRIIKTLDE		
SEQ ID NO: 11	SYSVRN WVAD	ATFV. HTNPG	L. .CDGCLAE	LGFWSSWKLV	RDDI IKELKE		
SEQ ID NO: 12	SSTIENWIAN	LDFILEDNDD	L. .CTGCKVH	TGFWKAWESA	ADELTSKIKS		
SEQ ID NO: 13	SSTIKNWIAD	LDFILQDND	L. .CTGCKVH	TGFWKAWESA	ADNLTSKIKS		
SEQ ID NO: 14	SADLANWVAN	LNFGLEDASD	L. .CSGCEVH	SGFWKAWSEI	ADTITSKVES		
	151					200	
SEQ ID NO: 1	AVREHPDYRV	VFTGHS LGGA	LATVAGADLR	GNGY.D	IDVFSYGAPR		175
SEQ ID NO: 2	ARKANPSFNV	ISTGHS LGGA	VAVLAAANLR	VGGT.P	VDIYTYGSPR		173
SEQ ID NO: 3	QLDRHPGYKI	VVTGHS LGGA	TAVLSALDLY	HHGHA.N	IEIYTQGQPR		
SEQ ID NO: 4	QLDRHPGYKI	VVTGHS LGGA	TAVLSALDLY	HHGHD.N	IEIYTQGQPR		
SEQ ID NO: 5	QFKQYPSYKV	AVTGHS LGGA	TALLCALDLY	QREEGLSSSN	LFLYTQGQPR		
SEQ ID NO: 6	QLTAHPTYKV	IVTGHS LGGA	QALLAGMDLY	QREPRLSPKN	LSIFTVGGPR		
SEQ ID NO: 7	QASQYPDYAL	TVTGHSLGAS	MAALTA AQL.	SATYD.N	VRLYTFGEPR		
SEQ ID NO: 8	QVSQFPDYAL	TVTGHSLGAS	LAALTA AQL.	SATYD.N	IRLYTFGEPR		
SEQ ID NO: 9	AKTANPTFKF	VVTGHS LGGA	VATIAAAYLR	KDGF.P	FDLYTYGSPR		
SEQ ID NO: 10	LKPEHSDYKI	VVVGHS LGAA	IASLAAADLR	TKNY.D	AILYAYAAPR		

Fig. 1 cont.

SEQ ID NO: 11 VVAQNPNYEL VVVGHSLGAA VATLAATDLR GKGYP....S AKLYAYASPR
 SEQ ID NO: 12 AMSTYSGYTL YFTGHSLGGA LATLGATVLR NDGY.....S VELYTYGCPR
 SEQ ID NO: 13 AMSTYSGYTL YFTGHSLGGA LATLGATVLR NDGY.....S VELYTYGCPR
 SEQ ID NO: 14 ALSDHSYSL VLTGHSYGAA LAALAATALR NSGH.....S VELYNYGQPR

201

SEQ ID NO: 1 VGNRAFAEFL TVQ.....T GGTLYRITHT NDIVPRLPPR EFGYSHSSPE 219
 SEQ ID NO: 2 VGNAQLSAFV SNQ..... AGGEYRVTHA DDPVPRLPPL IFGYRHTTPE 216
 SEQ ID NO: 3 IGTPAFANYV IGT..... KIPYQRLVHE RDIVPHLPPG AFGFLHAGEE
 SEQ ID NO: 4 IGTPEFANYV IGT..... KIPYQRLVNE RDIVPHLPPG AFGFLHAGEE
 SEQ ID NO: 5 VGDDPAFANYV VST..... GIPYRRTVNE RDIVPHLPPA AFGFLHAGEE
 SEQ ID NO: 6 VGNPTFAYYV EST..... GIPFQRTVHK RDIVPHVPPQ SFGFLHPGVE
 SEQ ID NO: 7 SGNQAFASYM NDAFQVSSPE TTQYFRVTHS NDGIPNLPPA DEGYAHGGVE
 SEQ ID NO: 8 S.NQAFASYM NDAFQASSPD TTQYFRVTHA NDGIPNLPPA DEGYAHGVVE
 SEQ ID NO: 9 VGNDFFANFV TQQ..... TGAEYRVTHG DDPVPRLPPI VFGYRHTSPE
 SEQ ID NO: 10 VANKPLAEFI TNQ..... .GNNYRFTHN DDPVPKLPPL TMGYVHISPE
 SEQ ID NO: 11 VGNAALAKYI TAQ..... .GNNFRFTHT NDPVPKLPPL SMGYVHVSPE
 SEQ ID NO: 12 IGNYALAEHI TSQ.....G SGANFRVTHL NDIVPRVPPM DFGFSQPSPE
 SEQ ID NO: 13 VGNYALAEHI TSQ.....G SGANFPVTHL NDIVPRVPPM DFGFSQPSPE
 SEQ ID NO: 14 LGNEALATYI TDQ.....N KGGNYRVTHT NDIVPKLPPT LLGYHHFSPE

251

SEQ ID NO: 1 YWIKS..GTL VPVTRNDIVK IEGIDATG.G NNQPNIP... DIPAPHLWYF. 262
 SEQ ID NO: 2 FWLSGGGGDK VDYTISDVKV CEGAANLG.C NGGTLGL... DIAAHLHYF. 261
 SEQ ID NO: 3 FWIMK..... .DSSLRV CPNGIETDNC SNSIVPFT.. SVIDHLSYLD
 SEQ ID NO: 4 FWIMK..... .DSSLRV CPNGIETDNC SNSIVPFT.. SVIDHLSYLD
 SEQ ID NO: 5 YWITD..... .NSPETVQV CTSDELETSDC SNSIVPFT.. SVLDHLSYFG
 SEQ ID NO: 6 SWIKS..... .GTSNVQI CTSEIETKDC SNSIVPFT.. SILDHLSYFD
 SEQ ID NO: 7 YWSV....DP YSAQNTFVCT GDEVQCCE.A QGGQGVN... .DAHTTYF.
 SEQ ID NO: 8 YWSV....DP YSAQNTFVCT GDEVQCCE.A QGGQGVN... .NAHTTYF.
 SEQ ID NO: 9 YWLNG.GPLD KDYTVTEIKV CEGIANVM.C NGGTIGL... DILAHITYF.
 SEQ ID NO: 10 YYITA..PDN TTVTDNQVTV LDGYVNF.K.G NTGTSGGLPD LLAHSHVWY
 SEQ ID NO: 11 YWITS..PNN ATVSTSDIKV IDGDVSFD.G NTGTGLPLLT DFEAHLWYF.
 SEQ ID NO: 12 YWITS..GNG ASVTASDIEV IEGINSTA.G NAGEATV... SVLAHLWYF.
 SEQ ID NO: 13 YWITS..GTG ASVTASDIEL IEGINSTA.G NAGEATV... DVLHLWYF.
 SEQ ID NO: 14 YYISS..ADE ATVTTTVDVTE VTGIDATG.G NDGTDGT... SIDAHRWYF.

301

SEQ ID NO: 1 GLIGT.CL.. 350
 SEQ ID NO: 2 QATDA.CNAG GFS..... 269
 SEQ ID NO: 3 MNTGL.CL.. 286
 SEQ ID NO: 4 MNTGL.CL..
 SEQ ID NO: 5 INTGL.CT..
 SEQ ID NO: 6 INEGS.CL..
 SEQ ID NO: 7 GMTSGACTW.
 SEQ ID NO: 8 GMTSGHCTW
 SEQ ID NO: 9 QSMAT.CAPI AIPWKR....
 SEQ ID NO: 10 FIHADACKGP GLPLR.....
 SEQ ID NO: 11 VQVDAGKGP LPFKR.....
 SEQ ID NO: 12 FAISE.CLL.
 SEQ ID NO: 13 FAISE.CLL.
 SEQ ID NO: 14 IYISE.CS..

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SEQUENCE LISTING

<110> Novozymes A/S

<120> Lipolytic enzyme variants

<130> 10470-WO

<160> 14

<170> PatentIn version 3.2

<210> 1

<211> 269

<212> PRT

<213> Thermomyces lanuginosus

<400> 1

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Ser Ala Ala Ala Tyr Cys Gly Lys Asn Asn Asp Ala Pro Ala Gly Thr
 20 25 30

Asn Ile Thr Cys Thr Gly Asn Ala Cys Pro Glu Val Glu Lys Ala Asp
 35 40 45

Ala Thr Phe Leu Tyr Ser Phe Glu Asp Ser Gly Val Gly Asp Val Thr
 50 55 60

Gly Phe Leu Ala Leu Asp Asn Thr Asn Lys Leu Ile Val Leu Ser Phe
 65 70 75 80

Arg Gly Ser Arg Ser Ile Glu Asn Trp Ile Gly Asn Leu Asn Phe Asp
 85 90 95

Leu Lys Glu Ile Asn Asp Ile Cys Ser Gly Cys Arg Gly His Asp Gly
 100 105 110

Phe Thr Ser Ser Trp Arg Ser Val Ala Asp Thr Leu Arg Gln Lys Val
 115 120 125

Glu Asp Ala Val Arg Glu His Pro Asp Tyr Arg Val Val Phe Thr Gly
 130 135 140

His Ser Leu Gly Gly Ala Leu Ala Thr Val Ala Gly Ala Asp Leu Arg
 145 150 155 160

Gly Asn Gly Tyr Asp Ile Asp Val Phe Ser Tyr Gly Ala Pro Arg Val
 165 170 175

Gly Asn Arg Ala Phe Ala Glu Phe Leu Thr Val Gln Thr Gly Gly Thr
 180 185 190

Leu Tyr Arg Ile Thr His Thr Asn Asp Ile Val Pro Arg Leu Pro Pro

195 200 10356-WO.ST25 205

Arg Glu Phe Gly Tyr Ser His Ser Ser Pro Glu Tyr Trp Ile Lys Ser
210 215 220

Gly Thr Leu Val Pro Val Thr Arg Asn Asp Ile Val Lys Ile Glu Gly
225 230 235 240

Ile Asp Ala Thr Gly Gly Asn Asn Gln Pro Asn Ile Pro Asp Ile Pro
245 250 255

Ala His Leu Trp Tyr Phe Gly Leu Ile Gly Thr Cys Leu
260 265

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<213> Fusarium oxysporum

<400> 2

Ala Val Gly Val Thr Thr Thr Asp Phe Ser Asn Phe Lys Phe Tyr Ile
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Gln His Gly Ala Ala Ala Tyr Cys Asn Ser Glu Ala Ala Ala Gly Ser
20 25 30

Lys Ile Thr Cys Ser Asn Asn Gly Cys Pro Thr Val Gln Gly Asn Gly
35 40 45

Ala Thr Ile Val Thr Ser Phe Val Gly Ser Lys Thr Gly Ile Gly Gly
50 55 60

Tyr Val Ala Thr Asp Ser Ala Arg Lys Glu Ile Val Val Ser Phe Arg
65 70 75 80

Gly Ser Ile Asn Ile Arg Asn Trp Leu Thr Asn Leu Asp Phe Gly Gln
85 90 95

Glu Asp Cys Ser Leu Val Ser Gly Cys Gly Val His Ser Gly Phe Gln
100 105 110

Arg Ala Trp Asn Glu Ile Ser Ser Gln Ala Thr Ala Ala Val Ala Ser
115 120 125

Ala Arg Lys Ala Asn Pro Ser Phe Asn Val Ile Ser Thr Gly His Ser
130 135 140

Leu Gly Gly Ala Val Ala Val Leu Ala Ala Ala Asn Leu Arg Val Gly
145 150 155 160

Gly Thr Pro Val Asp Ile Tyr Thr Tyr Gly Ser Pro Arg Val Gly Asn
165 170 175

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Ala Gln Leu Ser Ala Phe Val Ser Asn Gln Ala Gly Gly Glu Tyr Arg
180 185 190

Val Thr His Ala Asp Asp Pro Val Pro Arg Leu Pro Pro Leu Ile Phe
195 200 205

Gly Tyr Arg His Thr Thr Pro Glu Phe Trp Leu Ser Gly Gly Gly Gly
210 215 220

Asp Lys Val Asp Tyr Thr Ile Ser Asp Val Lys Val Cys Glu Gly Ala
225 230 235 240

Ala Asn Leu Gly Cys Asn Gly Gly Thr Leu Gly Leu Asp Ile Ala Ala
245 250 255

His Leu His Tyr Phe Gln Ala Thr Asp Ala Cys Asn Ala Gly Gly Phe
260 265 270

Ser Trp Arg Arg Tyr Arg Ser Ala Glu Ser Val Asp Lys Arg
275 280 285

<210> 3
<211> 265
<212> PRT
<213> Absidia reflexa

<400> 3

Ser Ser Ser Ser Thr Gln Asp Tyr Arg Ile Ala Ser Glu Ala Glu Ile
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Lys Ala His Thr Phe Tyr Thr Ala Leu Ser Ala Asn Ala Tyr Cys Arg
20 25 30

Thr Val Ile Pro Gly Gly Arg Trp Ser Cys Pro His Cys Gly Val Ala
35 40 45

Ser Asn Leu Gln Ile Thr Lys Thr Phe Ser Thr Leu Ile Thr Asp Thr
50 55 60

Asn Val Leu Val Ala Val Gly Glu Lys Glu Lys Thr Ile Tyr Val Val
65 70 75 80

Phe Arg Gly Thr Ser Ser Ile Arg Asn Ala Ile Ala Asp Ile Val Phe
85 90 95

Val Pro Val Asn Tyr Pro Pro Val Asn Gly Ala Lys Val His Lys Gly
100 105 110

Phe Leu Asp Ser Tyr Asn Glu Val Gln Asp Lys Leu Val Ala Glu Val
115 120 125

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Lys Ala Gln Leu Asp Arg His Pro Gly Tyr Lys Ile Val Val Thr Gly
130 135 140

His Ser Leu Gly Gly Ala Thr Ala Val Leu Ser Ala Leu Asp Leu Tyr
145 150 155 160

His His Gly His Ala Asn Ile Glu Ile Tyr Thr Gln Gly Gln Pro Arg
165 170 175

Ile Gly Thr Pro Ala Phe Ala Asn Tyr Val Ile Gly Thr Lys Ile Pro
180 185 190

Tyr Gln Arg Leu Val His Glu Arg Asp Ile Val Pro His Leu Pro Pro
195 200 205

Gly Ala Phe Gly Phe Leu His Ala Gly Glu Glu Phe Trp Ile Met Lys
210 215 220

Asp Ser Ser Leu Arg Val Cys Pro Asn Gly Ile Glu Thr Asp Asn Cys
225 230 235 240

Ser Asn Ser Ile Val Pro Phe Thr Ser Val Ile Asp His Leu Ser Tyr
245 250 255

Leu Asp Met Asn Thr Gly Leu Cys Leu
260 265

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<211> 264

<212> PRT

<213> Absidia corymbifera

<400> 4

Ser Ser Ser Thr Gln Asp Tyr Arg Ile Ala Ser Glu Ala Glu Ile Lys
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Ala His Thr Phe Tyr Thr Ala Leu Ser Ala Asn Ala Tyr Cys Arg Thr
20 25 30

Val Ile Pro Gly Gly Gln Trp Ser Cys Pro His Cys Asp Val Ala Pro
35 40 45

Asn Leu Asn Ile Thr Lys Thr Phe Thr Thr Leu Ile Thr Asp Thr Asn
50 55 60

Val Leu Val Ala Val Gly Glu Asn Glu Lys Thr Ile Tyr Val Val Phe
65 70 75 80

Arg Gly Thr Ser Ser Ile Arg Asn Ala Ile Ala Asp Ile Val Phe Val
85 90 95

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Pro Val Asn Tyr Pro Pro Val Asn Gly Ala Lys Val His Lys Gly Phe
 100 105 110

Leu Asp Ser Tyr Asn Glu Val Gln Asp Lys Leu Val Ala Glu Val Lys
 115 120 125

Ala Gln Leu Asp Arg His Pro Gly Tyr Lys Ile Val Val Thr Gly His
 130 135 140

Ser Leu Gly Gly Ala Thr Ala Val Leu Ser Ala Leu Asp Leu Tyr His
 145 150 155 160

His Gly His Asp Asn Ile Glu Ile Tyr Thr Gln Gly Gln Pro Arg Ile
 165 170 175

Gly Thr Pro Glu Phe Ala Asn Tyr Val Ile Gly Thr Lys Ile Pro Tyr
 180 185 190

Gln Arg Leu Val Asn Glu Arg Asp Ile Val Pro His Leu Pro Pro Gly
 195 200 205

Ala Phe Gly Phe Leu His Ala Gly Glu Glu Phe Trp Ile Met Lys Asp
 210 215 220

Ser Ser Leu Arg Val Cys Pro Asn Gly Ile Glu Thr Asp Asn Cys Ser
 225 230 235 240

Asn Ser Ile Val Pro Phe Thr Ser Val Ile Asp His Leu Ser Tyr Leu
 245 250 255

Asp Met Asn Thr Gly Leu Cys Leu
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 <212> PRT
 <213> Rhizomucor miehei

<400> 5

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Leu Thr Tyr Tyr Thr Thr Leu Ser Ala Asn Ser Tyr Cys Arg Thr Val
 20 25 30

Ile Pro Gly Ala Thr Trp Asp Cys Ile His Cys Asp Ala Thr Glu Asp
 35 40 45

Leu Lys Ile Ile Lys Thr Trp Ser Thr Leu Ile Tyr Asp Thr Asn Ala
 50 55 60

Met Val Ala Arg Gly Asp Ser Glu Lys Thr Ile Tyr Ile Val Phe Arg

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Val Pro Asp Gly Lys Ile Ile Thr Thr Phe Thr Ser Leu Leu Ser Asp
 50 55 60
 Thr Asn Gly Tyr Val Leu Arg Ser Asp Lys Gln Lys Thr Ile Tyr Leu
 65 70 75 80
 Val Phe Arg Gly Thr Asn Ser Phe Arg Ser Ala Ile Thr Asp Ile Val
 85 90 95
 Phe Asn Phe Ser Asp Tyr Lys Pro Val Lys Gly Ala Lys Val His Ala
 100 105 110
 Gly Phe Leu Ser Ser Tyr Glu Gln Val Val Asn Asp Tyr Phe Pro Val
 115 120 125
 Val Gln Glu Gln Leu Thr Ala His Pro Thr Tyr Lys Val Ile Val Thr
 130 135 140
 Gly His Ser Leu Gly Gly Ala Gln Ala Leu Leu Ala Gly Met Asp Leu
 145 150 155 160
 Tyr Gln Arg Glu Pro Arg Leu Ser Pro Lys Asn Leu Ser Ile Phe Thr
 165 170 175
 Val Gly Gly Pro Arg Val Gly Asn Pro Thr Phe Ala Tyr Tyr Val Glu
 180 185 190
 Ser Thr Gly Ile Pro Phe Gln Arg Thr Val His Lys Arg Asp Ile Val
 195 200 205
 Pro His Val Pro Pro Gln Ser Phe Gly Phe Leu His Pro Gly Val Glu
 210 215 220
 Ser Trp Ile Lys Ser Gly Thr Ser Asn Val Gln Ile Cys Thr Ser Glu
 225 230 235 240
 Ile Glu Thr Lys Asp Cys Ser Asn Ser Ile Val Pro Phe Thr Ser Ile
 245 250 255
 Leu Asp His Leu Ser Tyr Phe Asp Ile Asn Glu Gly Ser Cys Leu
 260 265 270

<210> 7
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 <212> PRT
 <213> Aspergillus niger

<400> 7

Thr Ala Gly Gln Ala Leu Ala Ala Ser Thr Gln Gly Ile Ser Glu Asp
 1 5 10 15

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Leu Tyr Asn Arg Leu Val Glu Met Ala Thr Ile Ser Gln Ala Ala Tyr
 20 25 30
 Ala Asp Leu Cys Asn Ile Pro Ser Thr Ile Ile Lys Gly Glu Lys Ile
 35 40 45
 Tyr Asn Ala Gln Thr Asp Ile Asn Gly Trp Ile Leu Arg Asp Asp Thr
 50 55 60
 Ser Lys Glu Ile Ile Thr Val Phe Arg Gly Thr Gly Ser Asp Thr Asn
 65 70 75 80
 Leu Gln Leu Asp Thr Asn Tyr Thr Leu Thr Pro Phe Asp Thr Leu Pro
 85 90 95
 Gln Cys Asn Asp Cys Glu Val His Gly Gly Tyr Tyr Ile Gly Trp Ile
 100 105 110
 Ser Val Gln Asp Gln Val Glu Ser Leu Val Lys Gln Gln Ala Ser Gln
 115 120 125
 Tyr Pro Asp Tyr Ala Leu Thr Val Thr Gly His Ser Leu Gly Ala Ser
 130 135 140
 Met Ala Ala Leu Thr Ala Ala Gln Leu Ser Ala Thr Tyr Asp Asn Val
 145 150 155 160
 Arg Leu Tyr Thr Phe Gly Glu Pro Arg Ser Gly Asn Gln Ala Phe Ala
 165 170 175
 Ser Tyr Met Asn Asp Ala Phe Gln Val Ser Ser Pro Glu Thr Thr Gln
 180 185 190
 Tyr Phe Arg Val Thr His Ser Asn Asp Gly Ile Pro Asn Leu Pro Pro
 195 200 205
 Ala Asp Glu Gly Tyr Ala His Gly Gly Val Glu Tyr Trp Ser Val Asp
 210 215 220
 Pro Tyr Ser Ala Gln Asn Thr Phe Val Cys Thr Gly Asp Glu Val Gln
 225 230 235 240
 Cys Cys Glu Ala Gln Gly Gly Gln Gly Val Asn Asp Ala His Thr Thr
 245 250 255
 Tyr Phe Gly Met Thr Ser Gly Ala Cys Thr Trp
 260 265

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 <211> 266
 <212> PRT

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<213> Aspergillus tubingensis

<400> 8

Thr Ala Gly His Ala Leu Ala Ala Ser Thr Gln Gly Ile Ser Glu Asp
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Leu Tyr Ser Arg Leu Val Glu Met Ala Thr Ile Ser Gln Ala Ala Tyr
 20 25 30

Ala Asp Leu Cys Asn Ile Pro Ser Thr Ile Ile Lys Gly Glu Lys Ile
 35 40 45

Tyr Asn Ser Gln Thr Asp Ile Asn Gly Trp Ile Leu Arg Asp Asp Ser
 50 55 60

Ser Lys Glu Ile Ile Thr Val Phe Arg Gly Thr Gly Ser Asp Thr Asn
 65 70 75 80

Leu Gln Leu Asp Thr Asn Tyr Thr Leu Thr Pro Phe Asp Thr Leu Pro
 85 90 95

Gln Cys Asn Ser Cys Glu Val His Gly Gly Tyr Tyr Ile Gly Trp Ile
 100 105 110

Ser Val Gln Asp Gln Val Glu Ser Leu Val Gln Gln Gln Val Ser Gln
 115 120 125

Phe Pro Asp Tyr Ala Leu Thr Val Thr Gly His Ser Leu Gly Ala Ser
 130 135 140

Leu Ala Ala Leu Thr Ala Ala Gln Leu Ser Ala Thr Tyr Asp Asn Ile
 145 150 155 160

Arg Leu Tyr Thr Phe Gly Glu Pro Arg Ser Asn Gln Ala Phe Ala Ser
 165 170 175

Tyr Met Asn Asp Ala Phe Gln Ala Ser Ser Pro Asp Thr Thr Gln Tyr
 180 185 190

Phe Arg Val Thr His Ala Asn Asp Gly Ile Pro Asn Leu Pro Pro Ala
 195 200 205

Asp Glu Gly Tyr Ala His Gly Val Val Glu Tyr Trp Ser Val Asp Pro
 210 215 220

Tyr Ser Ala Gln Asn Thr Phe Val Cys Thr Gly Asp Glu Val Gln Cys
 225 230 235 240

Cys Glu Ala Gln Gly Gly Gln Gly Val Asn Asn Ala His Thr Thr Tyr
 245 250 255

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 20 25 30

His Cys Ser Ala Gly Asn Cys Pro Asp Ile Glu Lys Asp Ala Ala Ile
 35 40 45

Val Val Gly Ser Val Val Gly Thr Lys Thr Gly Ile Gly Ala Tyr Val
 50 55 60

Ala Thr Asp Asn Ala Arg Lys Glu Ile Val Val Ser Val Arg Gly Ser
 65 70 75 80

Ile Asn Val Arg Asn Trp Ile Thr Asn Phe Asn Phe Gly Gln Lys Thr
 85 90 95

Cys Asp Leu Val Ala Gly Cys Gly Val His Thr Gly Phe Leu Asp Ala
 100 105 110

Trp Glu Glu Val Ala Ala Asn Val Lys Ala Ala Val Ser Ala Ala Lys
 115 120 125

Thr Ala Asn Pro Thr Phe Lys Phe Val Val Thr Gly His Ser Leu Gly
 130 135 140

Gly Ala Val Ala Thr Ile Ala Ala Ala Tyr Leu Arg Lys Asp Gly Phe
 145 150 155 160

Pro Phe Asp Leu Tyr Thr Tyr Gly Ser Pro Arg Val Gly Asn Asp Phe
 165 170 175

Phe Ala Asn Phe Val Thr Gln Gln Thr Gly Ala Glu Tyr Arg Val Thr
 180 185 190

His Gly Asp Asp Pro Val Pro Arg Leu Pro Pro Ile Val Phe Gly Tyr
 195 200 205

Arg His Thr Ser Pro Glu Tyr Trp Leu Asn Gly Gly Pro Leu Asp Lys
 210 215 220

Asp Tyr Thr Val Thr Glu Ile Lys Val Cys Glu Gly Ile Ala Asn Val

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235

225 230 240

Met Cys Asn Gly Gly Thr Ile Gly Leu Asp Ile Leu Ala His Ile Thr
245 250 255Tyr Phe Gln Ser Met Ala Thr Cys Ala Pro Ile Ala Ile Pro Trp Lys
260 265 270

Arg

<210> 10
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<212> PRT
<213> Aspergillus oryzae

<400> 10

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20 25 30Lys Leu Asn Cys Ser Val Gly Asn Cys Pro Asp Val Glu Ala Ala Gly
35 40 45Ser Thr Val Lys Leu Ser Phe Ser Asp Asp Thr Ile Thr Asp Thr Ala
50 55 60Gly Phe Val Ala Val Asp Asn Thr Asn Lys Ala Ile Val Val Ala Phe
65 70 75 80Arg Gly Ser Tyr Ser Ile Arg Asn Trp Val Thr Asp Ala Thr Phe Pro
85 90 95Gln Thr Asp Pro Gly Leu Cys Asp Gly Cys Lys Ala Glu Leu Gly Phe
100 105 110Trp Thr Ala Trp Lys Val Val Arg Asp Arg Ile Ile Lys Thr Leu Asp
115 120 125Glu Leu Lys Pro Glu His Ser Asp Tyr Lys Ile Val Val Val Gly His
130 135 140Ser Leu Gly Ala Ala Ile Ala Ser Leu Ala Ala Ala Asp Leu Arg Thr
145 150 155 160Lys Asn Tyr Asp Ala Ile Leu Tyr Ala Tyr Ala Ala Pro Arg Val Ala
165 170 175Asn Lys Pro Leu Ala Glu Phe Ile Thr Asn Gln Gly Asn Asn Tyr Arg
180 185 190

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Phe Thr His Asn Asp Asp Pro Val Pro Lys Leu Pro Leu Leu Thr Met
 195 200 205

Gly Tyr Val His Ile Ser Pro Glu Tyr Tyr Ile Thr Ala Pro Asp Asn
 210 215 220

Thr Thr Val Thr Asp Asn Gln Val Thr Val Leu Asp Gly Tyr Val Asn
 225 230 235 240

Phe Lys Gly Asn Thr Gly Thr Ser Gly Gly Leu Pro Asp Leu Leu Ala
 245 250 255

Phe His Ser His Val Trp Tyr Phe Ile His Ala Asp Ala Cys Lys Gly
 260 265 270

Pro Gly Leu Pro Leu Arg
 275

<210> 11
 <211> 278
 <212> PRT
 <213> Penicillium camemberti

<400> 11

Asp Val Ser Thr Ser Glu Leu Asp Gln Phe Glu Phe Trp Val Gln Tyr
 1 5 10 15

Ala Ala Ala Ser Tyr Tyr Glu Ala Asp Tyr Thr Ala Gln Val Gly Asp
 20 25 30

Lys Leu Ser Cys Ser Lys Gly Asn Cys Pro Glu Val Glu Ala Thr Gly
 35 40 45

Ala Thr Val Ser Tyr Asp Phe Ser Asp Ser Thr Ile Thr Asp Thr Ala
 50 55 60

Gly Tyr Ile Ala Val Asp His Thr Asn Ser Ala Val Val Leu Ala Phe
 65 70 75 80

Arg Gly Ser Tyr Ser Val Arg Asn Trp Val Ala Asp Ala Thr Phe Val
 85 90 95

His Thr Asn Pro Gly Leu Cys Asp Gly Cys Leu Ala Glu Leu Gly Phe
 100 105 110

Trp Ser Ser Trp Lys Leu Val Arg Asp Asp Ile Ile Lys Glu Leu Lys
 115 120 125

Glu Val Val Ala Gln Asn Pro Asn Tyr Glu Leu Val Val Val Gly His
 130 135 140

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Ser Leu Gly Ala Ala Val Ala Thr Leu Ala Ala Thr Asp Leu Arg Gly
145 150 155 160

Lys Gly Tyr Pro Ser Ala Lys Leu Tyr Ala Tyr Ala Ser Pro Arg Val
165 170 175

Gly Asn Ala Ala Leu Ala Lys Tyr Ile Thr Ala Gln Gly Asn Asn Phe
180 185 190

Arg Phe Thr His Thr Asn Asp Pro Val Pro Lys Leu Pro Leu Leu Ser
195 200 205

Met Gly Tyr Val His Val Ser Pro Glu Tyr Trp Ile Thr Ser Pro Asn
210 215 220

Asn Ala Thr Val Ser Thr Ser Asp Ile Lys Val Ile Asp Gly Asp Val
225 230 235 240

Ser Phe Asp Gly Asn Thr Gly Thr Gly Leu Pro Leu Leu Thr Asp Phe
245 250 255

Glu Ala His Ile Trp Tyr Phe Val Gln Val Asp Ala Gly Lys Gly Pro
260 265 270

Gly Leu Pro Phe Lys Arg
275

<210> 12
<211> 270
<212> PRT
<213> Aspergillus foetidus

<400> 12

Ser Val Ser Thr Ser Thr Leu Asp Glu Leu Gln Leu Phe Ala Gln Trp
1 5 10 15

Ser Ala Ala Ala Tyr Cys Ser Asn Asn Ile Asp Ser Lys Asp Ser Asn
20 25 30

Leu Thr Cys Thr Ala Asn Ala Cys Pro Ser Val Glu Glu Ala Ser Thr
35 40 45

Thr Met Leu Leu Glu Phe Asp Leu Thr Asn Asp Phe Gly Gly Thr Ala
50 55 60

Gly Phe Leu Ala Ala Asp Asn Thr Asn Lys Arg Leu Val Val Ala Phe
65 70 75 80

Arg Gly Ser Ser Thr Ile Glu Asn Trp Ile Ala Asn Leu Asp Phe Ile
85 90 95

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Leu Glu Asp Asn Asp Asp Leu Cys Thr Gly Cys Lys Val His Thr Gly
 100 105 110

Phe Trp Lys Ala Trp Glu Ser Ala Ala Asp Glu Leu Thr Ser Lys Ile
 115 120 125

Lys Ser Ala Met Ser Thr Tyr Ser Gly Tyr Thr Leu Tyr Phe Thr Gly
 130 135 140

His Ser Leu Gly Gly Ala Leu Ala Thr Leu Gly Ala Thr Val Leu Arg
 145 150 155 160

Asn Asp Gly Tyr Ser Val Glu Leu Tyr Thr Tyr Gly Cys Pro Arg Ile
 165 170 175

Gly Asn Tyr Ala Leu Ala Glu His Ile Thr Ser Gln Gly Ser Gly Ala
 180 185 190

Asn Phe Arg Val Thr His Leu Asn Asp Ile Val Pro Arg Val Pro Pro
 195 200 205

Met Asp Phe Gly Phe Ser Gln Pro Ser Pro Glu Tyr Trp Ile Thr Ser
 210 215 220

Gly Asn Gly Ala Ser Val Thr Ala Ser Asp Ile Glu Val Ile Glu Gly
 225 230 235 240

Ile Asn Ser Thr Ala Gly Asn Ala Gly Glu Ala Thr Val Ser Val Leu
 245 250 255

Ala His Leu Trp Tyr Phe Phe Ala Ile Ser Glu Cys Leu Leu
 260 265 270

<210> 13
 <211> 270
 <212> PRT
 <213> Aspergillus niger

<400> 13

Ser Val Ser Thr Ser Thr Leu Asp Glu Leu Gln Leu Phe Ser Gln Trp
 1 5 10 15

Ser Ala Ala Ala Tyr Cys Ser Asn Asn Ile Asp Ser Asp Asp Ser Asn
 20 25 30

Val Thr Cys Thr Ala Asp Ala Cys Pro Ser Val Glu Glu Ala Ser Thr
 35 40 45

Lys Met Leu Leu Glu Phe Asp Leu Thr Asn Asn Phe Gly Gly Thr Ala
 50 55 60

Gly Phe Leu Ala Ala Asp Asn Thr Asn Lys Arg Leu Val Val Ala Phe

Leu Thr Cys Ser Val Gly Asn Cys Pro Leu Val Glu Ala Ala Ser Thr
35 40 45

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Gln Ser Leu Asp Glu Phe Asn Glu Ser Ser Ser Tyr Gly Asn Pro Ala
 50 55 60
 Gly Tyr Leu Ala Ala Asp Glu Thr Asn Lys Leu Leu Val Leu Ser Phe
 65 70 75 80
 Arg Gly Ser Ala Asp Leu Ala Asn Trp Val Ala Asn Leu Asn Phe Gly
 85 90 95
 Leu Glu Asp Ala Ser Asp Leu Cys Ser Gly Cys Glu Val His Ser Gly
 100 105 110
 Phe Trp Lys Ala Trp Ser Glu Ile Ala Asp Thr Ile Thr Ser Lys Val
 115 120 125
 Glu Ser Ala Leu Ser Asp His Ser Asp Tyr Ser Leu Val Leu Thr Gly
 130 135 140
 His Ser Tyr Gly Ala Ala Leu Ala Ala Leu Ala Ala Thr Ala Leu Arg
 145 150 155 160
 Asn Ser Gly His Ser Val Glu Leu Tyr Asn Tyr Gly Gln Pro Arg Leu
 165 170 175
 Gly Asn Glu Ala Leu Ala Thr Tyr Ile Thr Asp Gln Asn Lys Gly Gly
 180 185 190
 Asn Tyr Arg Val Thr His Thr Asn Asp Ile Val Pro Lys Leu Pro Pro
 195 200 205
 Thr Leu Leu Gly Tyr His His Phe Ser Pro Glu Tyr Tyr Ile Ser Ser
 210 215 220
 Ala Asp Glu Ala Thr Val Thr Thr Thr Asp Val Thr Glu Val Thr Gly
 225 230 235 240
 Ile Asp Ala Thr Gly Gly Asn Asp Gly Thr Asp Gly Thr Ser Ile Asp
 245 250 255
 Ala His Arg Trp Tyr Phe Ile Tyr Ile Ser Glu Cys Ser
 260 265